SEQUENCE LISTING

<110> Garnaat, Carl W.
Lowe, Keith S.
Roth, Bradley A.

<120> ZmAxig1 Polynucleotides and Methods of Use

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<150> US 60/217,942
<151> 2000-07-13

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acgcacatga ccgcagtgcg cgcggggctg atcaagggaa agtgatcgg atg gag ctg    178
                                                    Met Glu Leu
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gag ctc ggg ctc gcg ccg ccg aac ccg cat cag ccg ctg gct gcc gcc      226
Glu Leu Gly Leu Ala Pro Pro Asn Pro His Gln Pro Leu Ala Ala Ala
     5                  10                  15

gcc gag ttc gtc ggt ctc ctc agc agc tcg gct ggc tcg tgc ggg aac      274
Ala Glu Phe Val Gly Leu Leu Ser Ser Ser Ala Gly Ser Cys Gly Asn
 20                  25                  30                  35

aag agg gtt ctc ggc gac gcg ttc ggg gcc gcc aag gcg gcc acg ctt      322
Lys Arg Val Leu Gly Asp Ala Phe Gly Ala Ala Lys Ala Ala Thr Leu
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ccg ctc ttc gtc tgc gag gat ggc gac gga ggc ggc ggc gac cgc gac      370
Pro Leu Phe Val Cys Glu Asp Gly Asp Gly Gly Gly Gly Asp Arg Asp
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cgc gac ggc gtc gtc gac cat gaa cag caa agc aac aat gta ccc agg      418
Arg Asp Gly Val Val Asp His Glu Gln Gln Ser Asn Asn Val Pro Arg
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Lys Lys Arg Leu Val Gly Trp Pro Pro Val Lys Cys Ala Arg Arg Arg
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agc tgc ggc ggc ggg tac gtg aag gtg aag ctg gaa ggg gtg ccc atc      514
Ser Cys Gly Gly Gly Tyr Val Lys Val Lys Leu Glu Gly Val Pro Ile
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ggg cgg aag gtg gac gtg tcc atc cac ggc tcg tac cag gag ctg ctc      562
Gly Arg Lys Val Asp Val Ser Ile His Gly Ser Tyr Gln Glu Leu Leu
                120                 125                 130

cgc acg ctc gag agc atg ttc cct tcg ggt aac caa caa gat cat gca      610
Arg Thr Leu Glu Ser Met Phe Pro Ser Gly Asn Gln Gln Asp His Ala
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gaa gac gag gtg gtg gtc tcg cac gag cgc cgc cgt cgc cat cct tat      658
Glu Asp Glu Val Val Val Ser His Glu Arg Arg Arg Arg His Pro Tyr
        150                 155                 160

gta gtc acc tac gag gac ggc gaa ggg gac tgg ttg ctc gtc gga gat      706
Val Val Thr Tyr Glu Asp Gly Glu Gly Asp Trp Leu Leu Val Gly Asp
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gat gtg ccg tgg gag gtc ttt gtc aag tca gtg aag cgg ctc aag ata      754
Asp Val Pro Trp Glu Val Phe Val Lys Ser Val Lys Arg Leu Lys Ile
180                 185                 190                 195

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Leu Ala  *
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Cys Gly Asn Lys Arg Val Leu Gly Asp Ala Phe Gly Ala Ala Lys Ala
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Ala Thr Leu Pro Leu Phe Val Cys Glu Asp Gly Asp Gly Gly Gly Gly
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Asp Arg Asp Arg Asp Gly Val Val Asp His Glu Gln Gln Ser Asn Asn
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Val Pro Arg Lys Lys Arg Leu Val Gly Trp Pro Pro Val Lys Cys Ala
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Val Pro Ile Gly Arg Lys Val Asp Val Ser Ile His Gly Ser Tyr Gln
        115                 120                 125
Glu Leu Leu Arg Thr Leu Glu Ser Met Phe Pro Ser Gly Asn Gln Gln
    130                 135                 140
Asp His Ala Glu Asp Glu Val Val Val Ser His Glu Arg Arg Arg Arg
145                 150                 155                 160
His Pro Tyr Val Val Thr Tyr Glu Asp Gly Glu Gly Asp Trp Leu Leu
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Val Gly Asp Asp Val Pro Trp Glu Val Phe Val Lys Ser Val Lys Arg
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Leu Lys Ile Leu Ala
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<213> Zea mays

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<212> DNA
<213> Zea mays

<400> 4

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<213> Zea mays

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<212> DNA
<213> Zea mays

<400> 6
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<210> 7
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<212> DNA
<213> Artificial Sequence

<220>
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<400> 7
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<210> 8
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<212> DNA
<213> Zea mays

<400> 8
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<210> 9
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<212> DNA
<213> Zea mays

<400> 9
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<210> 10
<211> 23
<212> DNA
<213> Zea mays

<400> 10
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<210> 11
<211> 36
<212> DNA
<213> Zea mays

<220>
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<210> 12
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<212> DNA
<213> Zea mays

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cgcgcggggc tgatcaaggg aaagtgatcg gatggagctg 100

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<212> DNA
<213> Zea mays

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<210> 14
<211> 22
<212> DNA
<213> Zea mays

<400> 14
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<210> 15
<211> 22
<212> DNA
<213> Zea mays

<400> 15
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<210> 16
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<213> Zea mays

<400> 16

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<210> 17
<211> 1433
<212> DNA
<213> Zea mays

<400> 17

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<211> 1173
<212> DNA
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35 40 45
Ala Asn Xaa Arg Cys Xaa Xaa Xaa Xaa Arg Lys Thr Xaa Xaa Xaa Glu
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Xaa
65